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       Yayon, Avner

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<141> 2004-06-17

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<151> 2003-06-17

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<151> 2003-07-30

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Ser Ala Ala Trp Asn Trp Ile Arg Gln Ser Pro Gly Arg Gly Leu Glu
        35   40   45

Trp Leu Gly Arg Thr Tyr Tyr Arg Ser Lys Trp Tyr Asn Asp Tyr Ala
    50   55   60

Val Ser Val Lys Ser Arg Ile Thr Ile Asn Pro Asp Thr Ser Lys Asn
65   70   75   80

Gln Phe Ser Leu Gln Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val
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Tyr Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
        35   40   45

Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Gln Lys Phe
    50   55   60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65   70   75   80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85   90   95

Ala Arg Asp Phe Leu Gly Tyr Glu Phe Asp Tyr Trp Gly Gln Gly Thr
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Leu Val Thr Val Ser Ser
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            20   25   30

Gly Val Gly Val Gly Trp Ile Arg Gln Pro Pro Gly Lys Ala Leu Glu
        35   40   45

Trp Leu Ala Leu Ile Asp Trp Asp Asp Asp Lys Tyr Tyr Ser Thr Ser
    50   55   60

Leu Lys Thr Arg Leu Thr Ile Ser Lys Asp Thr Ser Lys Asn Gln Val
65   70   75   80

PRO-028 PCT.ST25.txt

Val Leu Thr Met Thr Asn Met Asp Pro Val Asp Thr Ala Thr Tyr Tyr
              85   90   95

Cys Ala Arg Tyr His Ser Trp Tyr Glu Met Gly Tyr Tyr Gly Ser Thr
              100   105   110

Val Gly Tyr Met Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val
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Ser Ser
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              20   25   30

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              35   40   45

Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
    50   55   60

Gln Gly Arg Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr Met
65   70   75   80

Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala
              85   90   95

Arg Asp Asn Trp Phe Lys Pro Phe Ser Asp Val Trp Gly Gln Gly Thr
              100   105   110

Leu Val Thr Val Ser Ser
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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser Tyr
                20              25              30

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        35                  40                  45

Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
    50              55                  60

Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
65                  70                  75                  80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Arg Val Asn His Trp Thr Tyr Thr Phe Asp Tyr Trp Gly Gln Gly
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Thr Leu Val Thr Val Ser Ser
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Tyr Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
        35                  40                  45

Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Gln Lys Phe
    50              55                  60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65                  70                  75                  80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Arg Gly Tyr Trp Tyr Ala Tyr Phe Thr Tyr Ile Asn Tyr Gly Tyr
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Phe Asp Asn Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
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            20   25   30

Tyr Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
        35   40   45

Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Gln Lys Phe
    50   55   60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65   70   75   80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85   90   95

Ala Arg Thr Trp Gln Tyr Ser Tyr Phe Tyr Tyr Leu Asp Gly Gly Tyr
            100   105   110

Tyr Phe Asp Ile Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
            20   25   30

Tyr Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
        35   40   45

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Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Gln Lys Phe
    50   55   60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65   70   75   80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85   90   95

Ala Arg Asn Met Ala Tyr Thr Asn Tyr Gln Tyr Val Asn Met Pro His
            100   105   110

Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            115   120   125

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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
            20   25   30

Tyr Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
        35   40   45

Gly Trp Ile Asn Pro Asn Ser Gly Gly Thr Asn Tyr Ala Gln Lys Phe
    50   55   60

Gln Gly Arg Val Thr Met Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65   70   75   80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85   90   95

Ala Arg Ser Met Asn Ser Thr Met Tyr Trp Tyr Leu Arg Arg Val Leu
            100   105   110

Phe Asp His Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
            115   120   125

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Thr Ala Arg Ile Ser Cys Ser Gly Asp Ala Leu Gly Asp Lys Tyr Ala
            20   25   30

Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35   40   45

Asp Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50   55   60

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Glu
65   70   75   80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Gly Pro Asp Leu Trp
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Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Gln
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Thr Ala Arg Ile Ser Cys Ser Gly Asp Ala Leu Gly Asp Lys Tyr Ala
            20   25   30

Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35   40   45

Asp Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50   55   60

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Glu
65   70   75   80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Ser Ala Asp Tyr
                85   90   95

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            20   25   30

Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35   40   45

Asp Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50   55   60

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Glu
65   70   75   80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Phe Asp Phe Ala Val
                85   90   95

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Ser Asn Asn Lys Asn Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
        35   40   45

Pro Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val
    50   55   60

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Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr
65   70   75   80

Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln
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Tyr Asp Ser Ile Pro Tyr Thr Phe Gly Gln Gly Thr Lys Val Glu Ile
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Lys Arg Thr
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                20   25   30

Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
            35   40   45

Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Val Pro Ala Arg Phe Ser
        50   55   60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu
65   70   75   80

Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Met Ser Asn Tyr Pro
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    50              55              60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65              70              75              80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Asn Asn
                85              90              95

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Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
            35              40              45

Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Val Pro Ala Arg Phe Ser
    50              55              60

Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu
65              70              75              80

Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Thr Asn Asn Ala Pro
            85              90              95

Val Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr
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            20   25   30

Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35   40   45

Asp Asp Ser Asp Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50   55   60

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Glu
65   70   75   80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Tyr Phe Lys Leu Val
                85   90   95

Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Gln
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            20   25   30

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        35   40   45

Met Ile Tyr Asp Val Ser Asn Arg Pro Ser Gly Val Ser Asn Arg Phe
    50   55   60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65   70   75   80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Tyr Asp Met Tyr
                85   90   95

Asn Tyr Ile Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Gln
            100   105   110

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Ala Trp Asn Trp Ile Arg Gln Ser Pro Gly Arg Gly Leu Glu Trp Leu
                20                  25                  30

Gly Arg Thr Tyr Tyr Arg Ser Lys Trp Tyr Asn Asp Tyr Ala Val Ser
            35                  40                  45

Val Lys Ser Arg Ile Thr Ile Asn Pro Asp Thr Ser Lys Asn Gln Phe
        50                  55                  60

Ser Leu Gln Leu Asn Ser Val Thr Pro Glu Asp Thr Ala Val Tyr Tyr
65                  70                  75                  80

Cys Ala Arg Ser Tyr Tyr Pro Asp Phe Asp Tyr Trp Gly Gln Gly Thr
                85                  90                  95

Leu Val Thr Val Ser Ser Ala Gly Gly Gly Ser Gly Gly Gly Gly Ser
            100                 105                 110

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Asp Ile Glu Leu Thr Gln
            115                 120                 125

Pro Pro Ser Val Ser Val Ala Pro Gly Gln Thr Ala Arg Ile Ser Cys
    130                 135                 140

Ser Gly Asp Ala Leu Gly Asp Lys Tyr Ala Ser Trp Tyr Gln Gln Lys
145                 150                 155                 160

Pro Gly Gln Ala Pro Val Leu Val Ile Tyr Asp Asp Ser Asp Arg Pro
                165                 170                 175

Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala
            180                 185                 190

Thr Leu Thr Ile Ser Gly Thr Gln Ala Glu Asp Glu Ala Asp Tyr Tyr
            195                 200                 205

Cys Gln Ser Tyr Asp Gly Pro Asp Leu Trp Val Phe Gly Gly Gly Thr
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gcggaagacg aagcggatta ttattgccag agctatgacg gtcctgatct ttgggtgttt     660

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<400>   40
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PRO-028 PCT.ST25.txt

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<213>   ARTIFICIAL SEQUENCE

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PRO-028 PCT.ST25.txt

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PRO-028 PCT.ST25.txt

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PRO-028 PCT.ST25.txt

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aacgattatg cggtgagcgt gaaaagccgg attaccatca acccggatac ttcgaaaaac    240

cagtttagcc tgcaactgaa cagcgtgacc ccggaagata cggccgtgta ttattgcgcg    300

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cctgggcagg gtctcgagtg gatgggctgg attaacccga atagcggcgg cacgaactac    180

gcgcagaagt ttcagggccg ggtgaccatg acccgtgata ccagcattag caccgcgtat    240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgtgatttt    300

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gtgctgacta tgaccaacat ggacccggtg gatacggcca cctattattg cgcgcgttat    300

cattcttggt atgagatggg ttattatggt tctactgttg gttatatgtt tgattattgg    360

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PRO-028 PCT.ST25.txt

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cctgggcagg gtctcgagtg gatgggcggc attattccga tttttggcac ggcgaactac       180

gcgcagaagt ttcagggccg ggtgaccatt accgcggatg aaagcaccag caccgcgtat       240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgtgataat       300

tggtttaagc cttttttctga tgtttggggc caaggcaccc tggtgacggt tagctcagc        359
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gcgcagaagt ttcagggccg ggtgaccatt accgcggatg aaagcaccag caccgcgtat       240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgtgttaat       300

cattggactt atactttttga ttattggggc caaggcaccc tggtgacggt tagctcagc        359
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gcgcagaagt ttcagggccg ggtgaccatg acccgtgata ccagcattag caccgcgtat       240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgtggttat       300

tggtatgctt attttactta tattaattat ggttattttg ataattgggg ccaaggcacc       360

ctggtgacgg ttagctcagc                                                     380
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<210>  63
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<212>  DNA
<213>  ARTIFICIAL SEQUENCE

<220>
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PRO-028 PCT.ST25.txt

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gcgcagaagt ttcagggccg ggtgaccatg acccgtgata ccagcattag caccgcgtat   240

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accctggtga cggttagctc agc   383

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cctgggcagg gtctcgagtg gatgggctgg attaacccga atagcggcgg cacgaactac   180

gcgcagaagt ttcagggccg ggtgaccatg acccgtgata ccagcattag caccgcgtat   240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgtaatatg   300

gcttatacta attatcagta tgttaatatg cctcattttg attattgggg ccaaggcacc   360

ctggtgacgg ttagctcagc   380

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gcgcagaagt ttcagggccg ggtgaccatg acccgtgata ccagcattag caccgcgtat   240

atggaactga gcagcctgcg tagcgaagat acggccgtgt attattgcgc gcgttctatg   300

aattctacta tgtattggta tcttcgtcgt gttctttttg atcattgggg ccaaggcacc   360

ctggtgacgg ttagctcagc   380

<210>   66
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<212>   DNA
<213>   ARTIFICIAL SEQUENCE

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<220>
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caggcgccag ttctggtgat ttatgatgat tctgaccgtc cctcaggcat cccggaacgc   180

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gacgaagcgg attattattg ccagagctat gacggtcctg atctttgggt gtttggcggc   300

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caggcgccag ttctggtgat ttatgatgat tctgaccgtc cctcaggcat cccggaacgc   180

tttagcggat ccaacagcgg caacaccgcg accctgacca ttagcggcac tcaggcggaa   240

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gacgaagcgg attattattg ccagagctat gactttgatt ttgctgtgtt tggcggcggc   300

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tggtaccagc agaaaccagg tcagccgccg aaactattaa tttattgggc atccacccgt     180
gaaagcgggg tcccggatcg ttttagcggc tctggatccg gcactgattt taccctgacc     240
atttcgtccc tgcaagctga agacgtggcg gtgtattatt gccagcagta tgattctatt     300
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ccaggtcaag caccgcgtct attaatttat ggcgcgagca gccgtgcaac tggggtcccg     180
gcgcgtttta gcggctctgg atccggcacg gattttaccc tgaccattag cagcctggaa     240
cctgaagact ttgcgactta ttattgccag cagatgtcta attatcctga tacctttggc     300
cagggtacga agttgaaat  taaacgtacg                                       330
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catcccggga aggcgccgaa actgatgatt tatgatgtga gcaaccgtcc ctcaggcgtg     180
agcaaccgtt ttagcggatc caaaagcggc aacaccgcga gcctgaccat tagcggcctg     240
caagcggaag acgaagcgga ttattattgc cagagctatg acaataattc tgatgttgtg     300
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PRO-028 PCT.ST25.txt
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ccaggtcaag caccgcgtct attaatttat ggcgcgagca gccgtgcaac tggggtcccg      180

gcgcgtttta gcggctctgg atccggcacg gattttaccc tgaccattag cagcctggaa      240

cctgaagact ttgcgactta ttattgccag cagactaata atgctcctgt tacctttggc      300

cagggtacga aagttgaaat taaacgtacg                                       330


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gacgaagcgg attattattg ccagagctat gactatttta agcttgtgtt tggcggcggc      300

acgaagttaa ccgttcttgg ccag                                             324


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catcccggga aggcgccgaa actgatgatt tatgatgtga gcaaccgtcc ctcaggcgtg      180

agcaaccgtt ttagcggatc caaaagcggc aacaccgcga gcctgaccat tagcggcctg      240

caagcggaag acgaagcgga ttattattgc cagagctatg acatgtataa ttatattgtg      300

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<210>  75
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<213>  HOMO SAPIENS

<300>
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<309>  1991-08-01
<313>  (1)..(375)
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            20   25   30

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        35   40   45

Leu Val Phe Gly Ser Gly Asp Ala Val Glu Leu Ser Cys Pro Pro Pro
    50   55   60

Gly Gly Gly Pro Met Gly Pro Thr Val Trp Val Lys Asp Gly Thr Gly
65   70   75   80

Leu Val Pro Ser Glu Arg Val Leu Val Gly Pro Gln Arg Leu Gln Val
            85   90   95

Leu Asn Ala Ser His Glu Asp Ser Gly Ala Tyr Ser Cys Arg Gln Arg
            100   105   110

Leu Thr Gln Arg Val Leu Cys His Phe Ser Val Arg Val Thr Asp Ala
        115   120   125

Pro Ser Ser Gly Asp Asp Glu Asp Gly Glu Asp Glu Ala Glu Asp Thr
    130   135   140

Gly Val Asp Thr Gly Ala Pro Tyr Trp Thr Arg Pro Glu Arg Met Asp
145   150   155   160

Lys Lys Leu Leu Ala Val Pro Ala Ala Asn Thr Val Arg Phe Arg Cys
                165   170   175

Pro Ala Ala Gly Asn Pro Thr Pro Ser Ile Ser Trp Leu Lys Asn Gly
            180   185   190

Arg Glu Phe Arg Gly Glu His Arg Ile Gly Gly Ile Lys Leu Arg His
        195   200   205

Gln Gln Trp Ser Leu Val Met Glu Ser Val Val Pro Ser Asp Arg Gly
    210   215   220

Asn Tyr Thr Cys Val Val Glu Asn Lys Phe Gly Ser Ile Arg Gln Thr
225   230   235   240

Tyr Thr Leu Asp Val Leu Glu Arg Ser Pro His Arg Pro Ile Leu Gln
                245   250   255

Ala Gly Leu Pro Ala Asn Gln Thr Ala Val Leu Gly Ser Asp Val Glu
            260   265   270

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Phe His Cys Lys Val Tyr Ser Asp Ala Gln Pro His Ile Gln Trp Leu
        275             280             285

Lys His Val Glu Val Asn Gly Ser Lys Val Gly Pro Asp Gly Thr Pro
        290             295             300

Tyr Val Thr Val Leu Lys Val Ser Leu Glu Ser Asn Ala Ser Met Ser
305             310             315             320

Ser Asn Thr Pro Leu Val Arg Ile Ala Arg Leu Ser Ser Gly Glu Gly
            325             330             335

Pro Thr Leu Ala Asn Val Ser Glu Leu Glu Leu Pro Ala Asp Pro Lys
            340             345             350

Trp Glu Leu Ser Arg Ala Arg Leu Thr Leu Gly Lys Pro Leu Gly Glu
            355             360             365

Gly Cys Phe Gly Gln Val Val Met Ala Glu Ala Ile Gly Ile Asp Lys
        370             375             380

Asp Arg Ala Ala Lys Pro Val Thr Val Ala Val Lys Met Leu Lys Asp
385             390             395             400

Asp Ala Thr Asp Lys Asp Leu Ser Asp Leu Val Ser Glu Met Glu Met
            405             410             415

Met Lys Met Ile Gly Lys His Lys Asn Ile Ile Asn Leu Leu Gly Ala
            420             425             430

Cys Thr Gln Gly Gly Pro Leu Tyr Val Leu Val Glu Tyr Ala Ala Lys
            435             440             445

Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Asp
450             455             460

Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu Glu Gln Leu Thr Phe Lys
465             470             475             480

Asp Leu Val Ser Cys Ala Tyr Gln Val Ala Arg Gly Met Glu Tyr Leu
            485             490             495

Ala Ser Gln Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Val Leu
            500             505             510

Val Thr Glu Asp Asn Val Met Lys Ile Ala Asp Phe Gly Leu Ala Arg
        515             520             525

Asp Val His Asn Leu Asp Tyr Tyr Lys Lys Thr Thr Asn Gly Arg Leu
        530             535             540
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Pro Val Lys Trp Met Ala Pro Glu Ala Leu Phe Asp Arg Val Tyr Thr
545                 550             555             560

His Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe
                565             570             575

Thr Leu Gly Gly Ser Pro Tyr Pro Gly Ile Pro Val Glu Glu Leu Phe
            580             585             590

Lys Leu Leu Lys Glu Gly His Arg Met Asp Lys Pro Ala Asn Cys Thr
        595             600             605

His Asp Leu Tyr Met Ile Met Arg Glu Cys Trp His Ala Ala Pro Ser
    610             615             620

Gln Arg Pro Thr Phe Lys Gln Leu Val Glu Asp Leu Asp Arg Val Leu
625             630             635             640

Thr Val Thr Ser Thr Asp Glu Tyr Leu Asp Leu Ser Ala Pro Phe Glu
            645             650             655

Gln Tyr Ser Pro Gly Gly Gln Asp Thr Pro Ser Ser Ser Ser Ser Gly
            660             665             670

Asp Asp Ser Val Phe Ala His Asp Leu Leu Pro Pro Ala Pro Pro Ser
    675             680             685

Ser Gly Gly Ser Arg Thr
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<212>  DNA
<213>  HOMO SAPIENS

<300>
<308>  NCBI/AF2451114
<309>  2002-03-28
<313>  (40)..(2124)

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PRO-028 PCT.ST25.txt

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